

SEQ ID NO:	Clone	K _d , nM	K _i , nM
195	14N	0.4	2.0
199	25N	0.9	
191	4N	0.7	23.3
189	1N	0.5	16.7
205	36N	8.27	
203	34N	0.8	
207	42N	0.8	266.7
196	16N	1.4	
270	Consensus		

Summary Structure

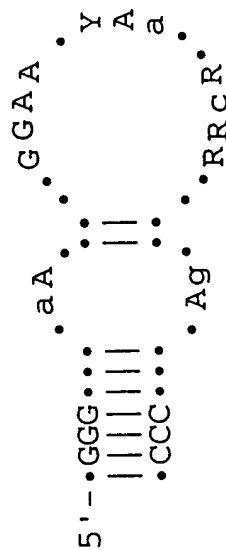


FIGURE 12C

